

104.6

**BIOTECH. LOGIC SYSTEMS BRANCH**

**RECEIVED**

**SCIENTIFIC INFORMATION CENTER**

**MAY 16 2001**

**1600/2900**

**RAW SEQUENCE LISTING  
ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/746,371A

Source: 1600

Date Processed by STIC: 5/10/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 3.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

**Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

**Checker Version 3.0 can be down loaded from the USPTO website at the following address:**  
<http://www.uspto.gov/web/offices/pac/checker>

# Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	<u>SERIAL NUMBER:</u> <u>09/946,371A</u>
<b>ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE</b>		
1 <input type="checkbox"/> Wrapped Nucleic	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3, as this will prevent "wrapping".	
2 <input type="checkbox"/> Wrapped Aminos	The amino acid number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3, as this will prevent "wrapping".	
3 <input type="checkbox"/> Incorrect Line Length	The rules require that a line not exceed 72 characters in length. This includes spaces.	
4 <input type="checkbox"/> Misaligned Amino Acid Numbering	The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.	
5 <input type="checkbox"/> Non-ASCII	This file was not saved in ASCII (DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text so that it can be processed.	
6 <input type="checkbox"/> Variable Length	Sequence(s) _____ contain n's or Xaa's which represented more than one residue. As per the rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.	
7 <input type="checkbox"/> PatentIn ver. 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
8 <input type="checkbox"/> Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please use the following format for each skipped sequence: <b>(2) INFORMATION FOR SEQ ID NO:X:</b> <b>(i) SEQUENCE CHARACTERISTICS:</b> (Do not insert any headings under "SEQUENCE CHARACTERISTICS") <b>(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:</b> This sequence is intentionally skipped  Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).	
9 <input type="checkbox"/> Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please use the following format for each skipped sequence. <210> sequence id number <400> sequence id number 000	
10 <input type="checkbox"/> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of <220> to <223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
11 <input type="checkbox"/> Use of "Artificial" (NEW RULES)	Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.	
12 <input type="checkbox"/> Use of <220>Feature (NEW RULES)	Sequence(s) _____ are missing the <220>Feature and associated headings. Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial Sequence" or "Unknown" Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)	
13 <input type="checkbox"/> PatentIn ver. 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other means to copy file to floppy disk.	

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/746,371A

DATE: 05/10/2001  
TIME: 12:21:36

Input Set : A:\BERL025.txt  
Output Set: N:\CRF3\05102001\I746371A.raw

*pyr 15*  
Does Not Comply  
Corrected Diskette Needed

3 <110> APPLICANT: Urry, Dan  
 5 <120> TITLE OF INVENTION: Acoustic Absorption Polymers and Their Methods of Use  
 7 <130> FILE REFERENCE: BERL025/01US  
 9 <140> CURRENT APPLICATION NUMBER: 09/746371A  
 10 <141> CURRENT FILING DATE: 2000-12-20  
 12 <160> NUMBER OF SEQ ID NOS: 47  
 14 <170> SOFTWARE: PatentIn version 3.0  
 16 <210> SEQ ID NO: 1  
 17 <211> LENGTH: 5  
 18 <212> TYPE: PRT  
 C--> 19 <213> ORGANISM: Artificial  
 21 <220> FEATURE:  
 22 <223> OTHER INFORMATION: This is a synthetic sequence.  
 24 <220> FEATURE:  
 25 <221> NAME/KEY: PEPTIDE  
 26 <222> LOCATION: (1)..(5)  
 28 <400> SEQUENCE: 1  
 30 Val Pro Gly Val Gly  
 31 1 5  
 33 <210> SEQ ID NO: 2  
 34 <211> LENGTH: 4  
 35 <212> TYPE: PRT  
 C--> 36 <213> ORGANISM: Artificial  
 38 <220> FEATURE:  
 39 <223> OTHER INFORMATION: This is a synthetic sequence.  
 41 <220> FEATURE:  
 42 <221> NAME/KEY: PEPTIDE  
 43 <222> LOCATION: (1)..(4)  
 45 <400> SEQUENCE: 2  
 47 Val Pro Gly Gly  
 48 1  
 50 <210> SEQ ID NO: 3  
 51 <211> LENGTH: 4  
 52 <212> TYPE: PRT  
 C--> 53 <213> ORGANISM: Artificial  
 55 <220> FEATURE:  
 56 <223> OTHER INFORMATION: This is a synthetic sequence.  
 58 <220> FEATURE:  
 59 <221> NAME/KEY: PEPTIDE  
 60 <222> LOCATION: (1)..(4)  
 62 <400> SEQUENCE: 3  
 64 Gly Gly Val Pro  
 65 1  
 67 <210> SEQ ID NO: 4  
 68 <211> LENGTH: 4  
 69 <212> TYPE: PRT  
 C--> 70 <213> ORGANISM: Artificial

RAW SEQUENCE LISTING  
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TIME: 12:21:36

Input Set : A:\BERL025.txt  
Output Set: N:\CRF3\05102001\I746371A.raw

72 <220> FEATURE:  
73 <223> OTHER INFORMATION: This is a synthetic sequence.

75 <220> FEATURE:  
76 <221> NAME/KEY: PEPTIDE  
77 <222> LOCATION: (1)..(4)  
79 <400> SEQUENCE: 4

81 Gly Gly Phe Pro

82 1

84 <210> SEQ ID NO: 5

85 <211> LENGTH: 4

86 <212> TYPE: PRT

C--> 87 <213> ORGANISM: Artificial

89 <220> FEATURE:

90 <223> OTHER INFORMATION: This is a synthetic sequence.

92 <220> FEATURE:

93 <221> NAME/KEY: PEPTIDE

94 <222> LOCATION: (1)..(4)

96 <400> SEQUENCE: 5

98 Gly Gly Ala Pro

99 1

101 <210> SEQ ID NO: 6

102 <211> LENGTH: 5

103 <212> TYPE: PRT

C--> 104 <213> ORGANISM: Artificial

106 <220> FEATURE:

107 <223> OTHER INFORMATION: This is a synthetic sequence.

109 <220> FEATURE:

110 <221> NAME/KEY: VARIANT

111 <222> LOCATION: (2)..(4)

112 <223> OTHER INFORMATION: Residue at position 2 is V, E, F, Y or K

113 Residue at position 4 is V, E, F or I

116 <400> SEQUENCE: 6

W--> 118 Gly Xaa Gly Xaa Pro

119 1 . . . 5

121 <210> SEQ ID NO: 7

122 <211> LENGTH: 6

123 <212> TYPE: PRT

C--> 124 <213> ORGANISM: Artificial

126 <220> FEATURE:

127 <223> OTHER INFORMATION: This is a synthetic sequence.

129 <220> FEATURE:

130 <221> NAME/KEY: PEPTIDE

131 <222> LOCATION: (1)..(6)

133 <400> SEQUENCE: 7

135 Ala Pro Gly Val Gly Val

136 1 . . . 5

138 <210> SEQ ID NO: 8

139 <211> LENGTH: 35

140 <212> TYPE: PRT

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/746,371A

DATE: 05/10/2001

TIME: 12:21:36

Input Set : A:\BERL025.txt

Output Set: N:\CRF3\05102001\I746371A.raw

C--&gt; 141 &lt;213&gt; ORGANISM: Artificial

143 &lt;220&gt; FEATURE:

144 &lt;223&gt; OTHER INFORMATION: This is a synthetic sequence.

146 &lt;220&gt; FEATURE:

147 &lt;221&gt; NAME/KEY: PEPTIDE

148 &lt;222&gt; LOCATION: (1)..(35)

150 &lt;400&gt; SEQUENCE: 8

152 Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Glu Gly Phe Pro Gly

153 1 5 10 15

155 Val Gly Val Pro Gly Val Gly Phe Pro Gly Phe Gly Phe Pro Gly Val

156 20 25 30

158 Gly Val Pro

159 35

161 &lt;210&gt; SEQ ID NO: 9

162 &lt;211&gt; LENGTH: 35

163 &lt;212&gt; TYPE: PRT

C--&gt; 164 &lt;213&gt; ORGANISM: Artificial

166 &lt;220&gt; FEATURE:

167 &lt;223&gt; OTHER INFORMATION: This is a synthetic sequence.

169 &lt;220&gt; FEATURE:

170 &lt;221&gt; NAME/KEY: PEPTIDE

171 &lt;222&gt; LOCATION: (1)..(35)

173 &lt;400&gt; SEQUENCE: 9

175 Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Glu Gly Phe Pro Gly

176 1 5 10 15

178 Val Gly Val Pro Gly Val Gly Phe Pro Gly Val Gly Phe Pro Gly Val

179 20 25 30

181 Gly Val Pro

182 35

184 &lt;210&gt; SEQ ID NO: 10

185 &lt;211&gt; LENGTH: 35

186 &lt;212&gt; TYPE: PRT

C--&gt; 187 &lt;213&gt; ORGANISM: Artificial

189 &lt;220&gt; FEATURE:

190 &lt;223&gt; OTHER INFORMATION: This is a synthetic sequence.

192 &lt;220&gt; FEATURE:

193 &lt;221&gt; NAME/KEY: PEPTIDE

194 &lt;222&gt; LOCATION: (1)..(35)

196 &lt;400&gt; SEQUENCE: 10

198 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Glu Gly Val Pro Gly

199 1 5 10 15

201 Val Gly Val Pro Gly Val Gly Phe Pro Gly Phe Gly Phe Pro Gly Val

202 20 25 30

204 Gly Val Pro

205 35

207 &lt;210&gt; SEQ ID NO: 11

208 &lt;211&gt; LENGTH: 35

209 &lt;212&gt; TYPE: PRT

C--&gt; 210 &lt;213&gt; ORGANISM: Artificial

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/746,371A

DATE: 05/10/2001  
TIME: 12:21:36

Input Set : A:\BERL025.txt  
Output Set: N:\CRF3\05102001\I746371A.raw

212 <220> FEATURE:  
213 <223> OTHER INFORMATION: This is a synthetic sequence.  
215 <220> FEATURE:  
216 <221> NAME/KEY: PEPTIDE  
217 <222> LOCATION: (1)..(35)  
219 <400> SEQUENCE: 11  
221 Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Glu Gly Phe Pro Gly  
222 1 5 10 15  
224 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val  
225 20 25 30  
227 Gly Val Pro  
228 35  
230 <210> SEQ ID NO: 12  
231 <211> LENGTH: 35  
232 <212> TYPE: PRT  
C--> 233 <213> ORGANISM: Artificial  
235 <220> FEATURE:  
236 <223> OTHER INFORMATION: This is a synthetic sequence.  
238 <220> FEATURE:  
239 <221> NAME/KEY: PEPTIDE  
240 <222> LOCATION: (1)..(35)  
242 <400> SEQUENCE: 12  
244 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Glu Gly Val Pro Gly  
245 1 5 10 15  
247 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val  
248 20 25 30  
250 Gly Val Pro  
251 35  
253 <210> SEQ ID NO: 13  
254 <211> LENGTH: 65  
255 <212> TYPE: PRT  
C--> 256 <213> ORGANISM: Artificial  
258 <220> FEATURE:  
259 <223> OTHER INFORMATION: This is a synthetic sequence.  
261 <220> FEATURE:  
262 <221> NAME/KEY: PEPTIDE  
263 <222> LOCATION: (1)..(65)  
265 <400> SEQUENCE: 13  
267 Gly Val Gly Ile Pro Gly Phe Gly Glu Pro Gly Glu Gly Phe Pro Gly  
268 1 5 10 15  
270 Val Gly Val Pro Gly Phe Gly Phe Pro Gly Phe Gly Ile Pro Gly Val  
271 20 25 30  
273 Gly Ile Pro Gly Phe Gly Glu Pro Gly Glu Gly Phe Pro Gly Val Gly  
274 35 40 45  
276 Val Pro Gly Phe Gly Phe Pro Gly Phe Gly Ile Pro Gly Val Gly Val  
277 50 55 60  
279 Pro  
280 65  
282 <210> SEQ ID NO: 14

RAW SEQUENCE LISTING DATE: 05/10/2001  
 PATENT APPLICATION: US/09/746,371A TIME: 12:21:36

Input Set : A:\BERL025.txt  
 Output Set: N:\CRF3\05102001\I746371A.raw

```

283 <211> LENGTH: 35
284 <212> TYPE: PRT
C--> 285 <213> ORGANISM: Artificial
287 <220> FEATURE:
288 <223> OTHER INFORMATION: This is a synthetic sequence.
290 <220> FEATURE:
291 <221> NAME/KEY: PEPTIDE
292 <222> LOCATION: (1)..(35)
294 <400> SEQUENCE: 14
296 Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Lys Gly Phe Pro Gly
297 1           5           10          15
299 Val Gly Val Pro Gly Val Gly Phe Pro Gly Phe Gly Phe Pro Gly Val
300           20          25          30
302 Gly Val Pro
303           35
305 <210> SEQ ID NO: 15
306 <211> LENGTH: 35
307 <212> TYPE: PRT
C--> 308 <213> ORGANISM: Artificial
310 <220> FEATURE:
311 <223> OTHER INFORMATION: This is a synthetic sequence.
313 <220> FEATURE:
314 <221> NAME/KEY: PEPTIDE
315 <222> LOCATION: (1)..(35)
317 <400> SEQUENCE: 15
319 Gly Val Gly Val Pro Gly Val Gly Phe Pro Gly Lys Gly Phe Pro Gly
320 1           5           10          15
322 Val Gly Val Pro Gly Val Gly Phe Pro Gly Val Gly Phe Pro Gly Val
323           20          25          30
325 Gly Val Pro
326           35
328 <210> SEQ ID NO: 16
329 <211> LENGTH: 35
330 <212> TYPE: PRT
C--> 331 <213> ORGANISM: Artificial
333 <220> FEATURE:
334 <223> OTHER INFORMATION: This is a synthetic sequence.
336 <220> FEATURE:
337 <221> NAME/KEY: PEPTIDE
338 <222> LOCATION: (1)..(35)
340 <400> SEQUENCE: 16
342 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Lys Gly Val Pro Gly
343 1           5           10          15
345 Val Gly Val Pro Gly Val Gly Phe Pro Gly Phe Gly Phe Pro Gly Val
346           20          25          30
348 Gly Val Pro
349           35
351 <210> SEQ ID NO: 17
352 <211> LENGTH: 35

```

*FJ*  
 The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

*FJ*  
**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields for each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/746,371A

DATE: 05/10/2001  
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Input Set : A:\\BERL025.txt  
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L:19 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1  
L:36 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2  
L:53 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3  
L:70 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4  
L:87 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5  
L:104 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6  
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:124 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7  
L:141 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8  
L:164 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9  
L:187 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10  
L:210 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11  
L:233 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12  
L:256 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13  
L:285 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14  
L:308 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15  
L:331 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16  
L:354 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17  
L:377 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:18  
L:400 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:19  
L:423 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:20  
L:446 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:21  
L:460 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21  
L:466 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:22  
L:483 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:23  
L:496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23  
L:502 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24  
L:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24  
L:523 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25  
L:543 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26  
L:561 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:27  
L:579 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:28  
L:596 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29  
L:619 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:30  
L:639 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:31  
L:659 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:32  
L:682 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:33  
L:702 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:34  
L:722 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:35  
L:757 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:36  
L:794 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:37  
L:820 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:38  
L:848 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:39  
L:871 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:40  
L:894 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:41  
L:914 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:42  
L:936 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:43  
L:953 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:44

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/746,371A

DATE: 05/10/2001  
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Input Set : A:\BERL025.txt  
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L:972 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:45  
L:989 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:46  
L:1008 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:47